

AMENDMENTSIn the claims

Please cancel Claims 15, 17, 18, 19, 20 and 24, without prejudice.

Please amend Claims 1 and 16 as follows:

Sub D1  
C1

Claim 1 (Twice Amended). A fabric softening protein hybrid comprising an amino acid sequence comprising a cellulose binding domain linked to a fabric softening protein; wherein said fabric softening protein is linked to said amino acid sequence comprising a cellulose binding domain, via an amino acid and/or non-amino acid linking region; wherein the cellulose binding domain is selected from the group consisting of CBD Cellulozome from *Clostridium cellulovorans*, CBD E3 from *Thermonospora fusca*, CBD-dimer from *Clostridium stercorarium* XynA, CBD from *Bacillus agaradherens*, and/or mixtures thereof; wherein 2 to 50 amino acid sequences are cross-linked via a non-amino acid linking region; and further wherein said non-amino linking region is a polymer selected from polyethylene glycol derivatives, nucleophilic polyethylene glycol derivatives, carboxyl polyethylene glycol derivatives, electrophilically activated polyethylene glycol derivatives, sulfhydryl-selective polyethylene glycol derivatives, heterofunctional polyethylene glycol derivatives, biotin polyethylene glycol derivatives, vinyl polyethylene glycol derivatives, silane polyethylene glycol derivatives, phospholipid polyethylene glycol derivatives, 1-ethyl-3-(3-dimethylaminopropyl) carbodiimide, N-ethyl-5-phenylisaoxolium-3-sulphonate, 1-cyclohexyl-3(2-morpholineethyl) carbodide metho-p-toluene sulphonate, N-ethoxycarbonyl-2-ethoxy 1,2 dihydroquinoline or glutaraldehyde and mixtures thereof.

Sub D2  
C2  
Sub D2

Claim 16 (Twice Amended). A fabric softening protein hybrid according to claim 1 wherein the amino acid sequence comprising a cellulose binding domain is selected from CBD Cellulozome from *Clostridium cellulovorans*.